## SEQUENCE LISTING

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tcggc	gatt a	tg ga et As 1 tcc	ac co sp Pr	eg go ro Al	cc ga La Gi aga	ag go lu Ai 5 gaa	cg g la Va gac	tg c: al Lo	tg ca eu Gi	aa galn G	ag ag lu Ly 10	ag go ys A	ca ct la Le	tc aag eu Lys cct	111
ttt at Phe Me 15	gatt a <sup>.</sup> Mo tg aat	tg ga et As 1 tcc Ser ata	tca Ser	gag Glu 20	ec ga La Gi aga Arg	ag go lu A 5 gaa Glu aat	cg g la Va gac Asp	tg c al L tgt Cys	tg ca eu G aat Asn 25	aa galn G	ag ag lu Ly 10 ggc Gly aca	ag goys A. gaa Glu	ca ct la Le ccc Pro	tc aag eu Lys cct Pro 30	111
ttt at Phe Me 15 agg aa Arg Ly	gatt a Mo tg aat et Asn ag ata	tg gaet As  1  tcc Ser  ata Ile	tca Ser cca Pro 35	gag Glu 20 gag Glu	aga Arg aag Lys	ag go lu A 5 gaa Glu aat Asn	gac Asp tca Ser	tgt Cys ctt Leu 40 aca	tg ca eu G aat Asn 25 aga Arg	aa galn Gaat Asn cag	ag ag lu Ly 10 ggc Gly aca Thr	gaa Glu tac Tyr	ca ci la Le ccc Pro aac Asn 45	tc aag eu Lys cct Pro 30 agc Ser	111
ttt at Phe Me 15 agg at Arg Ly	gatt a Mo tg aat et Asn ag ata ys Ile	tg gaet As  1 tcc Ser ata Ile ctc Leu 50 act	tca Ser cca Pro 35 tgc Cys	gag Glu 20 gag Glu tta Leu	aga Arg aag Lys aac Asn	ag go lu A 5 gaa Glu aat Asn caa Gln	gac Asp tca Ser gaa Glu 55	tgt Cys ctt Leu 40 aca Thr	aat Asn 25 aga Arg gta Val	aa galn Galn Cag Gln tgt Cys	ag ag lu Ly 10 ggc Gly aca Thr tta Leu	gaa Glu tac Tyr gca Ala 60	ca ctla Lecco Pro  aac Asn 45 agc Ser	tc aageu Lys  cct Pro 30  agc Ser  act Thr	111 159 207
ttt at Phe Mo 15 agg at Arg Ly tgt gc Cys Al Ala Mo act tc Thr Se	gatt a Mo	tg gaet As  1 tcc Ser ata Ile ctc Leu 50 act Thr	tca Ser cca Pro 35 tgc Cys	gag Glu 20 gag Glu tta Leu aat Asn	aga Arg aag Lys aac Asn tgt Cys	ag golu As Glu aat Asn Caa Gln 70	gac Asp tca Ser gaa Glu 55 gcc Ala	tgt Cys ctt Leu 40 aca Thr aaa	aat Asn 25 aga Arg gta Val aca Thr	aa galn Gaat Asn cag Gln tgt Cys aaa Lys	ag adlu Lyllo Color Colo	gaa Glu tac Tyr gca Ala 60 gcc Ala	ca ctla Le ccc Pro aac Asn 45 agc Ser aat Asn	cct Pro 30 agc Ser act Thr ggc Gly	111 159 207 255

					ttt Phe											447
					ata Ile											495
gat Asp	ttc Phe	ata Ile 145	act Thr	gct Ala	ctg Leu	cca Pro	gct Ala 150	cgg Arg	gga Gly	ttg Leu	gat Asp	cat His 155	atc Ile	gct Ala	gag Glu	543
aac Asn	att Ile 160	ctg Leu	tca Ser	tac Tyr	ctg Leu	gat Asp 165	gcc Ala	aaa Lys	tca Ser	cta Leu	tgt Cys 170	gct Ala	gct Ala	gaa Glu	ctt Leu	591
gtg Val 175	tgc Cys	aag Lys	gaa Glu	tgg Trp	tac Tyr 180	cga Arg	gtg Val	acc Thr	tct Ser	gat Asp 185	ggc Gly	atg Met	ctg Leu	tgg Trp	aag Lys 190	639
					atg Met											687
gca Ala	g <b>aa</b> Glu	cga Arg	aga Arg 210	gga Gly	tgg Trp	gga Gly	cag Gln	tat Tyr 215	tta Leu	ttc Phe	aaa Lys	aac Asn	aaa Lys 220	cct Pro	cct Pro	735
			-		ccc Pro					_	_					783
att Ile	ata Ile 240	caa Gln	gac Asp	att Ile	gag Glu	aca Thr 245	ata Ile	gaa Glu	tct Ser	aat Asn	tgg Trp 250	aga Arg	tgt Cys	gga Gly	aga Arg	831
cat His 255	agt Ser	tta Leu	cag Gln	aga Arg	att Ile 260	cac His	tgc Cys	cga Arg	agt Ser	gaa Glu 265	aca Thr	agc Ser	aaa Lys	gga Gly	gtt Val 270	879
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gtg Val	atc Ile 320	ata Ile	aca Thr	gga Gly	tca Ser	tcg Ser 325	gat Asp	tcc Ser	acg Thr	gtc Val	aga Arg 330	gtg Val	tgg Trp	gat Asp	gta Val	1071
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. 10.3

cgt tcc att Arg Ser Ile							1215
cgg agg gtg Arg Arg Val 385							1263
gat gac aag Asp Asp Lys 400							1311
tgg aac aca Trp Asn Thr 415		Glu Phe	Val Arg				1359
cga ggc att Arg Gly Ile							1407
tca tct gac Ser Ser Asp		Arg Leu					1455
tta cga gtg Leu Arg Val 465	tta gaa ggc Leu Glu Gly	cat gag His Glu 470	gaa ttg Glu Leu	gtg cgt Val Arg	tgt att Cys Ile 475	cga ttt Arg Phe	1503
gat aac aag Asp Asn Lys 480							1551
tgg gat ctt Trp Asp Leu 495			Pro Arg				1599
tgt cta cgg Cys Leu Arg							1647
ttt gat gaa Phe Asp Glu		Val Ser					1695
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tcc cct tct Ser Pro Ser 560	cga aca tac Arg Thr Tyr	acc tac Thr Tyr 565	atc tcc . Ile Ser .	aga taa Arg	ataacca t	acactgacc	1796
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cacaactgac t	gcttcagtg c	tgctatcag	aagatgt	ctt cta	tcaattg t	gaatgattg	2036
gaacttttaa a	cctcccctc c	tctcctcct	ttcacct	ctg cac	ctagttt t	ttcccattg	2096
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<213> Artificial sequence

<220>

<223> Description of the artificial sequence : ADNc coding for human  $\beta TrCP$  protein

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1 5 10 15

Asn Ser Ser Glu Arg Glu Asp Cys Asn Asn Gly Glu Pro Pro Arg Lys
20 25 30

Ile Ile Pro Glu Lys Asn Ser Leu Arg Gln Thr Tyr Asn Ser Cys Ala
35 40 45

Arg Leu Cys Leu Asn Gln Glu Thr Val Cys Leu Ala Ser Thr Ala Met 50 55 60

Lys Thr Glu Asn Cys Val Ala Lys Thr Lys Leu Ala Asn Gly Thr Ser 65 70 75 80

Ser Met Ile Val Pro Lys Gln Arg Lys Leu Ser Ala Ser Tyr Glu Lys 85 90 95

Glu Lys Glu Leu Cys Val Lys Tyr Phe Glu Gln Trp Ser Glu Ser Asp 100 105 110

Gln Val Glu Phe Val Glu His Leu Ile Ser Gln Met Cys His Tyr Gln 115 120 125

His Gly His Ile Asn Ser Tyr Leu Lys Pro Met Leu Gln Arg Asp Phe 130 135 140

Ile Thr Ala Leu Pro Ala Arg Gly Leu Asp His Ile Ala Glu Asn Ile 145 150 155 160

Leu Ser Tyr Leu Asp Ala Lys Ser Leu Cys Ala Ala Glu Leu Val Cys 165 170 175

Lys Glu Trp Tyr Arg Val Thr Ser Asp Gly Met Leu Trp Lys Lys Leu 180 185 190

Ile Glu Arg Met Val Arg Thr Asp Ser Leu Trp Arg Gly Leu Ala Glu 195 200 205

Arg Arg Gly Trp Gly Gln Tyr Leu Phe Lys Asn Lys Pro Pro Asp Gly 210 215 220

Asn Ala Pro Pro Asn Ser Phe Tyr Arg Ala Leu Tyr Pro Lys Ile Ile 225 230 235 240

Gln Asp Ile Glu Thr Ile Glu Ser Asn Trp Arg Cys Gly Arg His Ser 245 250 255

Leu Gln Arg Ile His Cys Arg Ser Glu Thr Ser Lys Gly Val Tyr Cys 260 265 270

Leu Gln Tyr Asp Asp Gln Lys Ile Val Ser Gly Leu Arg Asp Asn Thr 275 280 285

Ile Lys Ile Trp Asp Lys Asn Thr Leu Glu Cys Lys Arg Ile Leu Thr 290 295 300

Gly His Thr Gly Ser Val Leu Cys Leu Gln Tyr Asp Glu Arg Val Ile Ile Thr Gly Ser Ser Asp Ser Thr Val Arg Val Trp Asp Val Asn Thr Gly Glu Met Leu Asn Thr Leu Ile His His Cys Glu Ala Val Leu His 345 Leu Arg Phe Asn Asn Gly Met Met Val Thr Cys Ser Lys Asp Arg Ser 360 Ile Ala Val Trp Asp Met Ala Ser Pro Thr Asp Ile Thr Leu Arg Arg Val Leu Val Gly His Arg Ala Ala Val Asn Val Val Asp Phe Asp Asp Lys Tyr Ile Val Ser Ala Ser Gly Asp Arg Thr Ile Lys Val Trp Asn Thr Ser Thr Cys Glu Phe Val Arg Thr Leu Asn Gly His Lys Arg Gly Ile Ala Cys Leu Gln Tyr Arg Asp Arg Leu Val Val Ser Gly Ser Ser Asp Asn Thr Ile Arg Leu Trp Asp Ile Glu Cys Gly Ala Cys Leu Arg Val Leu Glu Gly His Glu Glu Leu Val Arg Cys Ile Arg Phe Asp Asn 470 Lys Arg Ile Val Ser Gly Ala Tyr Asp Gly Lys Ile Lys Val Trp Asp Leu Val Ala Ala Leu Asp Pro Arg Ala Pro Ala Gly Thr Leu Cys Leu Arg Thr Leu Val Glu His Ser Gly Arg Val Phe Arg Leu Gln Phe Asp 520 Glu Phe Gln Ile Val Ser Ser Ser His Asp Asp Thr Ile Leu Ile Trp Asp Phe Leu Asn Asp Pro Ala Ala Gln Ala Glu Pro Pro Arg Ser Pro 550 555 Ser Arg Thr Tyr Thr Tyr Ile Ser Arg

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<211> 19

<212> ADN

<213> Artificial sequence

565

<220>

<223> Description of the artificial sequence : sense primer

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<220> <223> Description of the artificial sequence : antisense primer	
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<210> 5 <211> 20 <212> ADN <213> Artificial sequence	
<220> <223> Description of the artificial sequence : sense primer	
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<220> <223> Description of the artificial sequence : antisense primer	
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<210> 8 <211> 29 <212> ADN <213> Artificial sequence	
<220> <223> Description of the artificial sequence : primer	
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